

RAW SEQUENCE LISTING

EFS

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/820,953A
Source: IFW/b
Date Processed by STIC: 4/5/07

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 04/05/2007

PATENT APPLICATION: US/10/820,953A

TIME: 15:53:41

Input Set : N:\EFS\04_05_07\10820953a_efs\sequence.txt

Output Set: N:\CRF4\04052007\J820953A.raw

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3 <110> APPLICANT: Koelsch, Gerald
4     Tang, Jordan J. N.
5     Hong, Lin
6     Ghosh, Arun K.
7     The Board of Trustees of the University of Illinois
8     Oklahoma Medical Research Foundation
10 <120> TITLE OF INVENTION: Inhibitors of Memapsin 2 and Use Thereof
12 <130> FILE REFERENCE: 022266-000930US
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/820,953A
C--> 15 <141> CURRENT FILING DATE: 2004-04-08
17 <150> PRIOR APPLICATION NUMBER: US 60/141,363
18 <151> PRIOR FILING DATE: 1999-06-28
20 <150> PRIOR APPLICATION NUMBER: US 60/168,060
21 <151> PRIOR FILING DATE: 1999-11-30
23 <150> PRIOR APPLICATION NUMBER: US 60/177,836
24 <151> PRIOR FILING DATE: 2000-01-25
26 <150> PRIOR APPLICATION NUMBER: US 60/178,368
27 <151> PRIOR FILING DATE: 2000-01-27
29 <150> PRIOR APPLICATION NUMBER: US 60/210,292
30 <151> PRIOR FILING DATE: 2000-06-08
32 <150> PRIOR APPLICATION NUMBER: US 09/603,713
33 <151> PRIOR FILING DATE: 2000-06-27
35 <150> PRIOR APPLICATION NUMBER: US 09/845,226
36 <151> PRIOR FILING DATE: 2001-04-30
38 <160> NUMBER OF SEQ ID NOS: 39
40 <170> SOFTWARE: PatentIn Ver. 2.1
42 <210> SEQ ID NO: 1
43 <211> LENGTH: 3252
44 <212> TYPE: DNA
45 <213> ORGANISM: Homo sapiens
47 <220> FEATURE:
48 <223> OTHER INFORMATION: memapsin 2
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58 gtcactgtgc gtgccaacat tgctgccatc actgaatcag acaagttctt catcaacggc 480
59 tccaactggg aaggcatcct ggggctggcc tatgctgaga ttgccaggcc tgacgactcc 540
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63 atccggcggg agtgggtatta tgagggtgatc attgtgcggg tggagatcaa tggacaggat 780
64 ctgaaaatgg actgcaagga gtacaactat gacaagagca ttgtggacag tggcaccacc 840
65 aaccttcgtt tgcccaagaa agtgtttgaa gctgcagtca aatccatcaa ggcagcctcc 900
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68 aaccagtcct tccgcatcac catccttccg cagcaatacc tgcggccagt ggaagatgtg 1080
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103 acctgataag ggagaggga atacaaggag ggcctctggg gttcctggcc tcagccagct 3180
104 gccmcaagc cataaacaa taaamcaaga atactgagtc taaaaaaaaa aaaaaaaaaa 3240
105 aaaaaaaaaa aa 3252
108 <210> SEQ ID NO: 2
109 <211> LENGTH: 488
110 <212> TYPE: PRT
111 <213> ORGANISM: Homo sapiens

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113 <220> FEATURE:
114 <223> OTHER INFORMATION: purified memapsin 2, aspartic proteinase 2 (ASP2)
116 <220> FEATURE:
117 <223> OTHER INFORMATION: amino acids 28-48 are remnant putative propeptide
118     residues
122 <220> FEATURE:
123 <223> OTHER INFORMATION: amino acids 54-57, 61-68, 73-80, 86-89, 109-111,
124     113-118, 123-134, 143-154, 165-168, 198-202, and
125     220-224 are N-lobe beta strands
127 <220> FEATURE:
128 <223> OTHER INFORMATION: amino acids 58-61, 78, 80, 82-83, 116, 118-121,
129     156, 166, 174, 246, 274, 276, 278-281, 283, and
130     376-377 are residues in contact with the OM99-2
131     inhibitor
133 <220> FEATURE:
134 <223> OTHER INFORMATION: amino acids 184-191 and 210-217 are N-lobe helices
136 <220> FEATURE:
137 <223> OTHER INFORMATION: amino acids 237-240, 247-249, 251-256, 259-260,
138     273-275, 282-285, 316-318, 331-336, 342-348,
139     354-357, 366-370, 372-375, 380-383, 390-395,
140     400-405, and 418-420 are C-lobe beta strands
142 <220> FEATURE:
143 <223> OTHER INFORMATION: amino acids 286-299, 307-310, 350-353, 384-387,
144     and 427-431 are C-lobe helices
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151           20           25           30
153 Glu Thr Asp Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val
154           35           40           45
156 Glu Met Val Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val
157           50           55           60
159 Glu Met Thr Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp
160   65           70           75           80
162 Thr Gly Ser Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu
163           85           90           95
165 His Arg Tyr Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg
166           100          105          110
168 Lys Gly Val Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu
169           115          120          125
171 Gly Thr Asp Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg
172           130          135          140
174 Ala Asn Ile Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly
175   145          150          155          160
177 Ser Asn Trp Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg
178           165          170          175
180 Pro Asp Asp Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr
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184 His Val Pro Asn Leu Phe Ser Leu Gln Leu Cys Gly Ala Gly Phe Pro
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187 Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile
188      210      215      220
190 Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro
191 225      230      235      240
193 Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile
194      245      250      255
196 Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys
197      260      265      270
199 Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val
200      275      280      285
202 Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys
203      290      295      300
205 Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala
206 305      310      315      320
208 Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met
209      325      330      335
211 Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln
212      340      345      350
214 Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr
215      355      360      365
217 Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val
218      370      375      380
220 Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile
221 385      390      395      400
223 Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala
224      405      410      415
226 Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr
227      420      425      430
229 Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val
230      435      440      445
232 Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val
233      450      455      460
235 Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe
236 465      470      475      480
238 Ala Asp Asp Ile Ser Leu Leu Lys
239      485
243 <210> SEQ ID NO: 3
244 <211> LENGTH: 503
245 <212> TYPE: PRT
246 <213> ORGANISM: Homo sapiens
248 <220> FEATURE:
249 <223> OTHER INFORMATION: pro-memapsin 2
251 <220> FEATURE:
252 <223> OTHER INFORMATION: amino acids 1-13 are the T7 promoter
254 <220> FEATURE:
255 <223> OTHER INFORMATION: amino acids 1-15 are vector-derived residues
257 <220> FEATURE:

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/05/2007
PATENT APPLICATION: US/10/820,953A TIME: 15:53:42

Input Set : N:\EFS\04_05_07\10820953a_efs\sequence.txt
Output Set: N:\CRF4\04052007\J820953A.raw

FyI

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:8; Xaa Pos. 2
Seq#:22; Xaa Pos. 3
Seq#:23; Xaa Pos. 1
Seq#:27; Xaa Pos. 3,4
Seq#:29; Xaa Pos. 2,3
Seq#:35; Xaa Pos. 4,5

VERIFICATION SUMMARY

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Input Set : N:\EFS\04_05_07\10820953a_efs\sequence.txt

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L:14 M:270 C: Current Application Number differs, Replaced Current Application Number
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:633 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:717 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:789 M:259 W: Allowed number of lines exceeded, <223> Other Information:
L:794 M:283 W: Missing Blank Line separator, <220> field identifier
L:916 M:283 W: Missing Blank Line separator, <220> field identifier
L:929 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0